

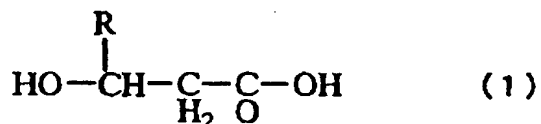
CLAIMS

1. A transformant

wherein at least one kind of gene expression cassette
5 comprising a polyester synthesis-associated enzyme gene has been
introduced into a yeast.

2. The transformant according to Claim 1

wherein the polyester is a copolymer resulting from the
10 copolymerization of 3-hydroxyalkanoic acids of the following
general formula (1);

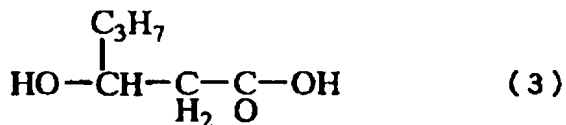
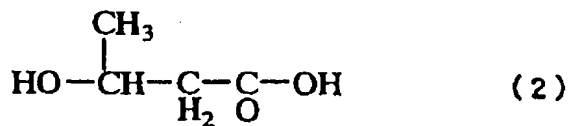


in the formula, R represents an alkyl group.

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3. The transformant according to Claim 1 or 2

wherein the polyester is copolyester P(3HB-co-3HH)
resulting from the copolymerization of 3-hydroxybutyric acid
of the following formula (2) and 3-hydroxyhexanoic acid of the
20 following formula (3);



4. The transformant according to any of Claims 1 to 3

wherein the yeast belongs to any of the genera
25 Aciculoconidium, Ambrosiozyma, Arthroascus, Arxiozyma, Ashbya,
Babjevia, Bensingtonia, Botryoascus, Botryozyma, Brettanomyces,

- Bullera, Bulleromyces, Candida, Citeromyces, Clavispora,
Cryptococcus, Cystofilobasidium, Debaryomyces, Dekkara,
Dipodascopsis, Dipodascus, Eeniella, Endomycopsella, Eremascus,
Eremothecium, Erythrobasidium, Fellomyces, Filobasidium,
5 Galactomyces, Geotrichum, Guilliermondella, Hanseniaspora,
Hansenula, Hasegawaea, Holtermannia, Hormoascus, Hyphopichia,
Issatchenkia, Kloeckera, Kloeckeraspora, Kluyveromyces, Kondoa,
Kuraishia, Kurtzmanomyces, Leucosporidium, Lipomyces,
Lodderomyces, Malassezia, Metschnikowia, Mrakia, Myxozyma,
10 Nadsonia, Nakazawaea, Nematospora, Ogataea, Oosporidium,
Pachysolen, Phachytichospora, Phaffia, Pichia, Rhodosporidium,
Rhodotorula, Saccharomyces, Saccharomycodes, Saccharomycopsis,
Saitoella, Sakaguchia, Saturnospora, Schizoblastosporion,
Schizosaccharomyces, Schwanniomyces, Sporidiobolus,
15 Sporobolomyces, Sporopachydermia, Stephanoascus,
Sterigmatomyces, Sterigmatosporidium, Symbiotaphrina,
Sympodiomyces, Sympodiomycopsis, Torulaspora, Trichosporiella,
Trichosporon, Trigonopsis, Tsuchiyaea, Udeniomyces, Waltomyces,
Wickerhamia, Wickerhamiella, Williopsis, Yamadazyma, Yarrowia,
20 Zygoascus, Zygosaccharomyces, Zygowilliopsis and Zygozyma.

5. The transformant according to any of Claims 1 to 4
wherein the yeast is Yarrowia lipolytica.
- 25 6. The transformant according to any of Claims 1 to 4
wherein the yeast is Candida maltosa.
7. The transformant according to any of Claims 1 to 6
wherein a polyester synthesis-associated enzyme gene
30 expression cassette comprises a promoter and a terminator,
said promoter and said terminator functioning in a yeast.
8. The transformant according to Claim 7
wherein the promoter and terminator are derived from
35 Yarrowia lipolytica.

9. The transformant according to Claim 7 or 8
wherein the promoter is derived from Yarrowia lipolytica
ALK3.

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10. The transformant according to Claim 7 or 8
wherein the terminator is derived from Yarrowia
lipolytica XPR2.

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11. The transformant according to Claim 7
wherein the promoter and terminator are derived from
Candida maltosa.

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12. The transformant according to Claim 7 or 11
wherein the promoter is derived from Candida maltosa ALK1.

13. The transformant according to Claim 7 or 11
wherein the terminator is derived from Candida maltosa
ALK1.

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14. The transformant according to any of Claims 1 to 13
wherein the polyester synthesis-associated enzyme gene
is derived from Aeromonas caviae.

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15. The transformant according to any of Claims 1 to 13
wherein the polyester synthesis-associated enzyme gene
is a PHA synthase gene derived from Aeromonas caviae or a PHA
synthase gene and (R)-specific enoyl-CoA hydratase gene.

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16. The transformant according to Claim 15
wherein said PHA synthase gene has the sequence represented
by SEQ ID NO:3

and the (R)-specific enoyl-CoA hydratase gene has the
sequence represented by SEQ ID NO:4.

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17. A method of producing a polyester using the transformant according to any of Claims 1 to 16 which comprises growing said transformant and harvesting a polyester from the resulting culture.

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18. A polyester synthesis-associated enzyme gene which is modified from at least one gene code CTG to TTA, TTG, CTT, CTC or CTA.

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19. The polyester synthesis-associated enzyme gene according to Claim 18 which codes for an enzyme derived from a bacterium.

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20. The polyester synthesis-associated enzyme gene according to Claim 19 wherein said bacterium is Aeromonas caviae.

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21. The polyester synthesis-associated enzyme gene according to Claim 20 wherein the enzyme gene derived from Aeromonas caviae is a PHA synthase gene or a (R)-specific enoyl-CoA hydratase gene.

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22. The polyester synthesis-associated enzyme gene according to Claim 21 wherein said PHA synthase gene has the sequence represented by SEQ ID NO:3.

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23. The polyester synthesis-associated enzyme gene according to Claim 21 wherein said (R)-specific enoyl-CoA hydratase gene has the sequence represented by SEQ ID NO:4.